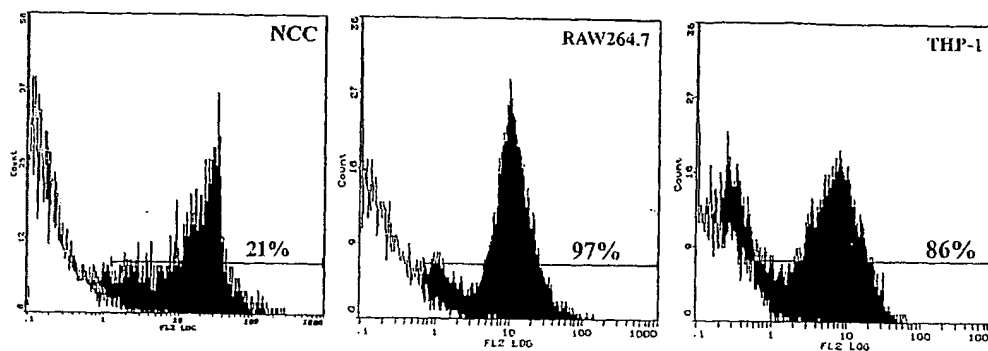


## Figures 1A - 1B

A



B

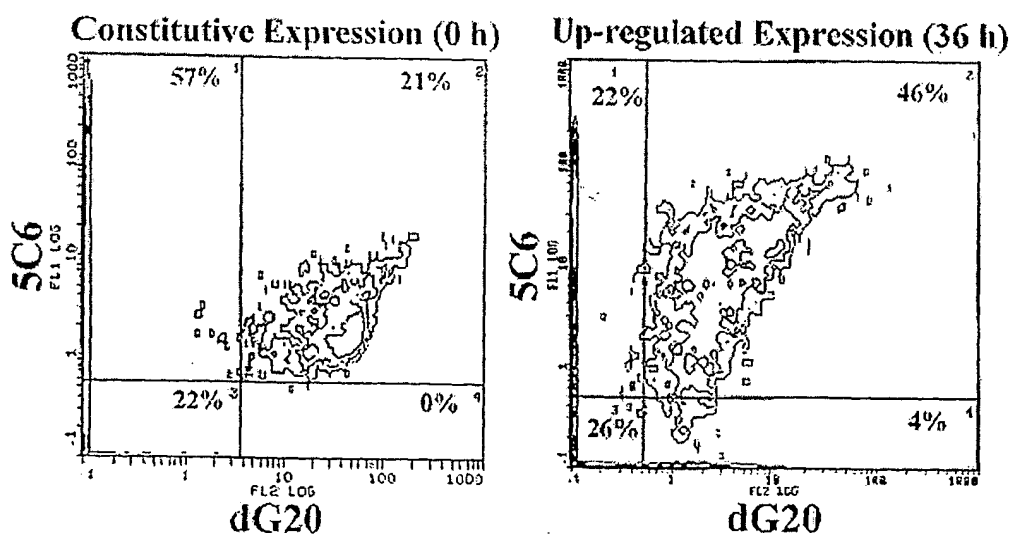
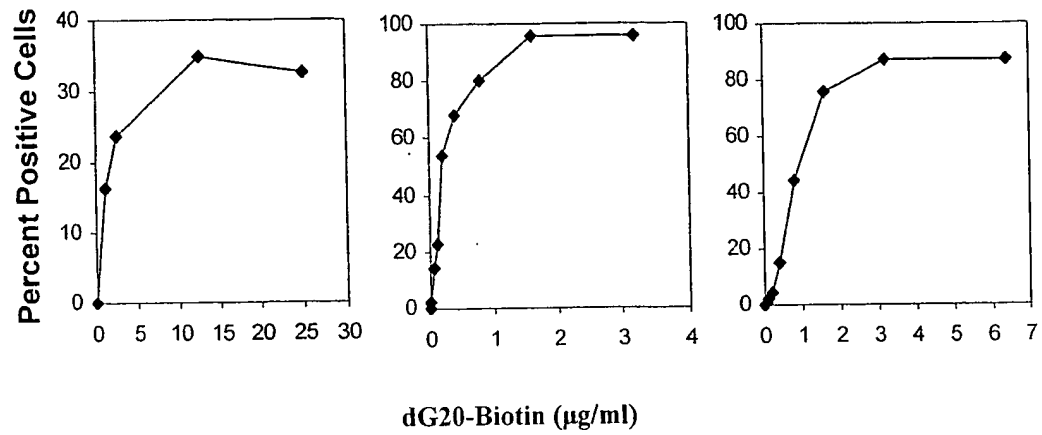
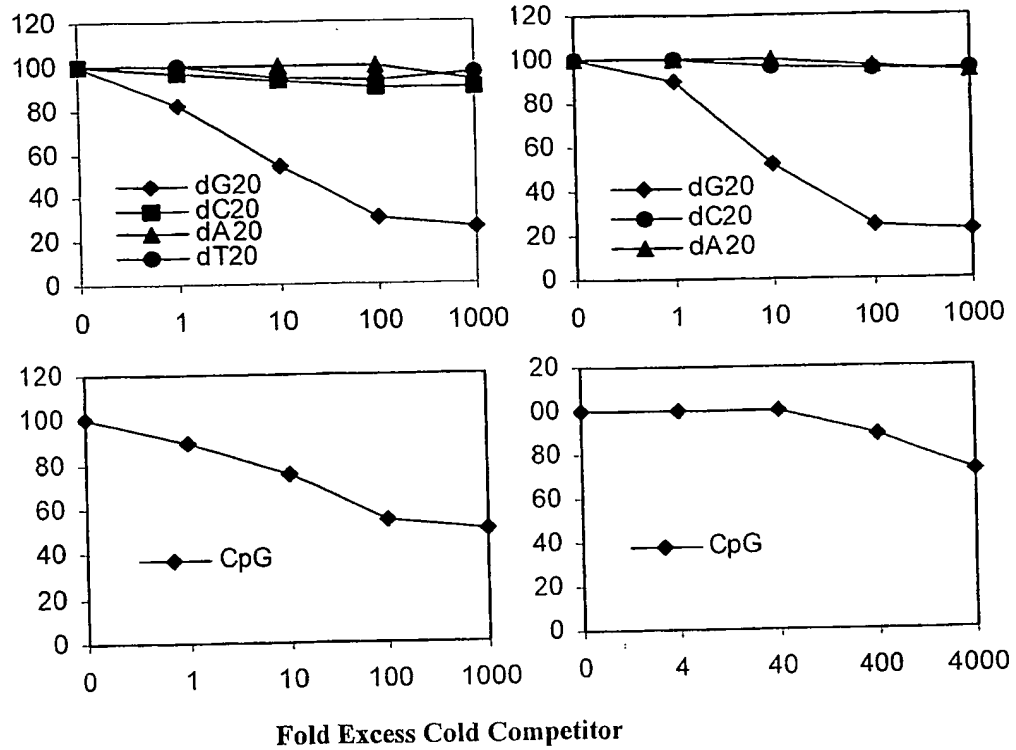


Figure 2

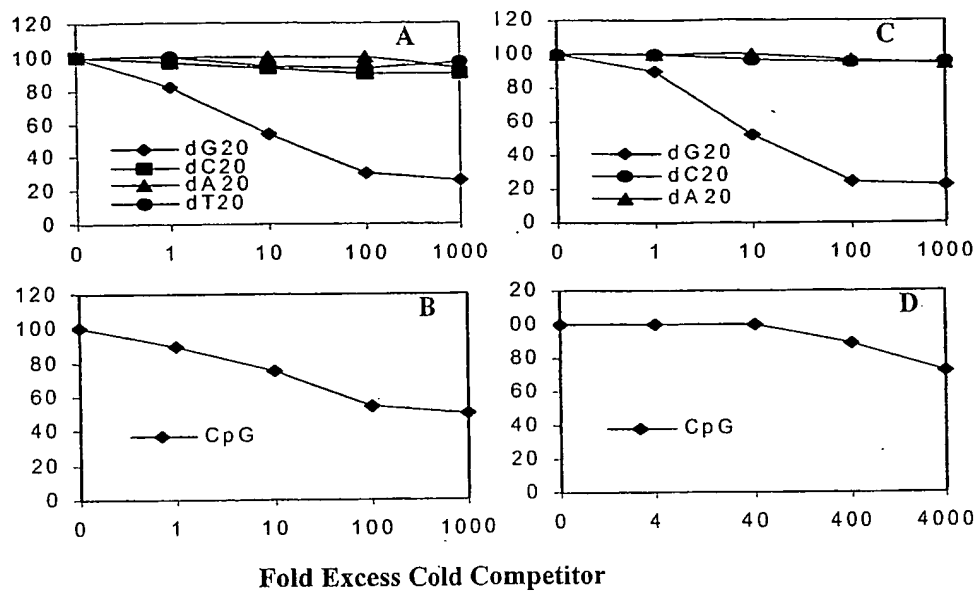


Figures 3A - 3D

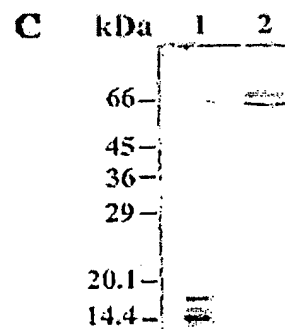
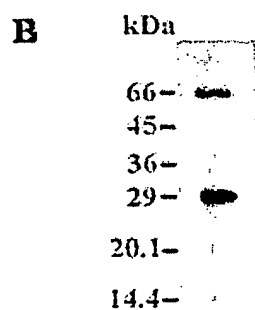
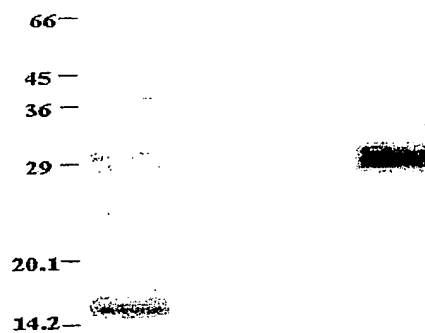


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Figures 4A - 4C



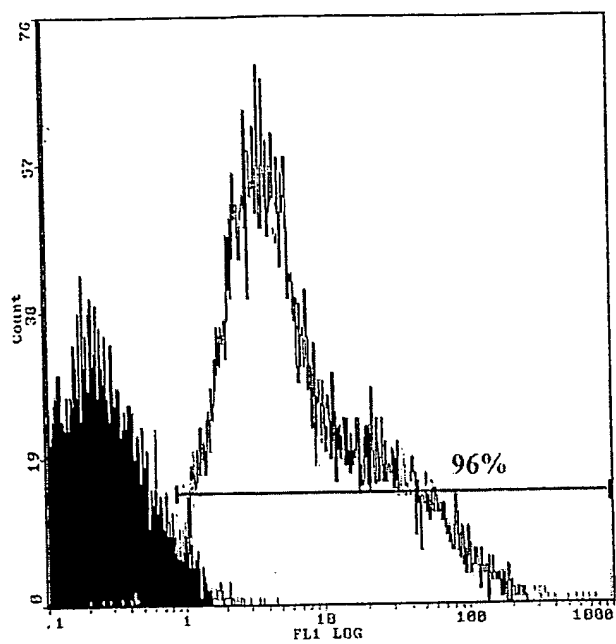
**A** 1 2 3 4 5 6



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Figures 5A - 5B

A



B

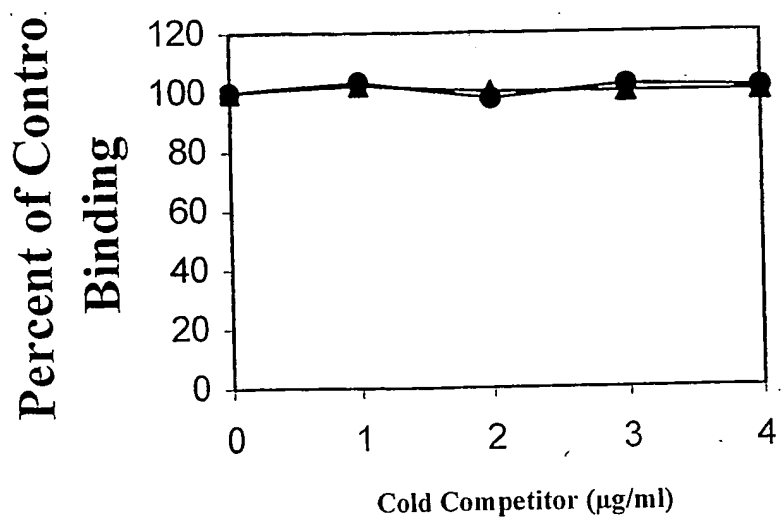
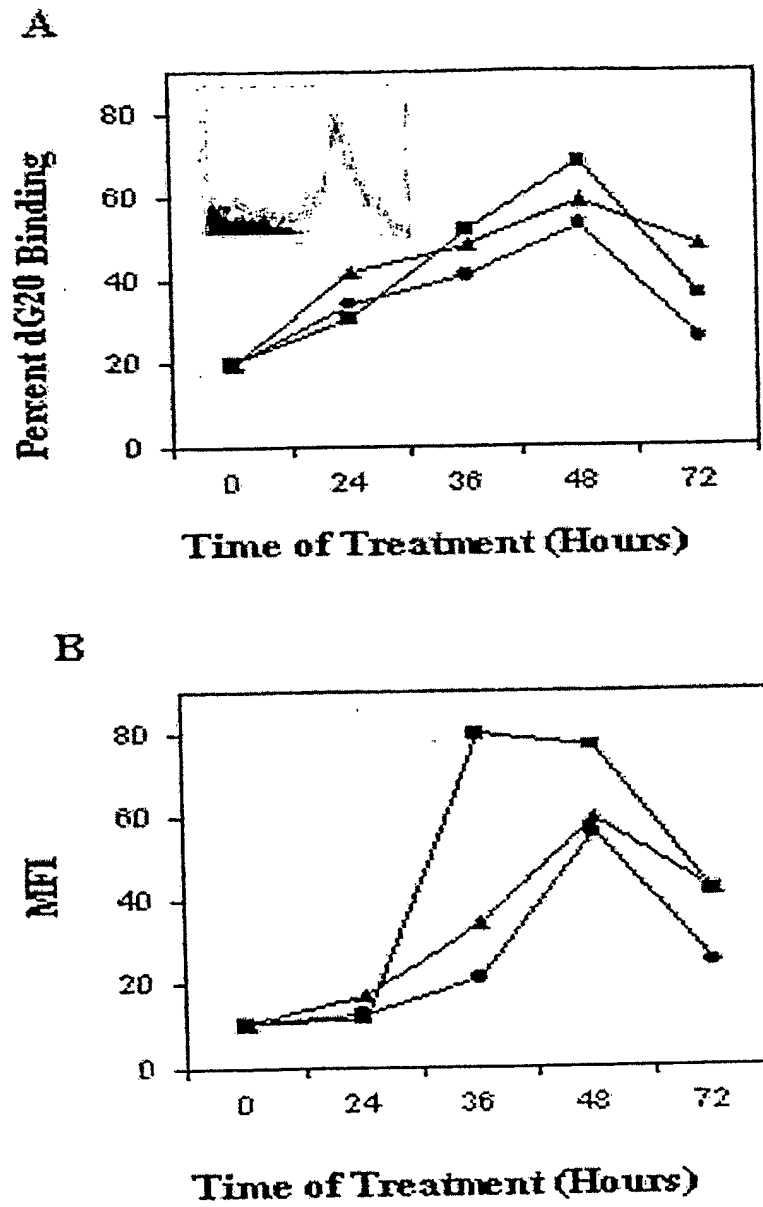


Figure 6



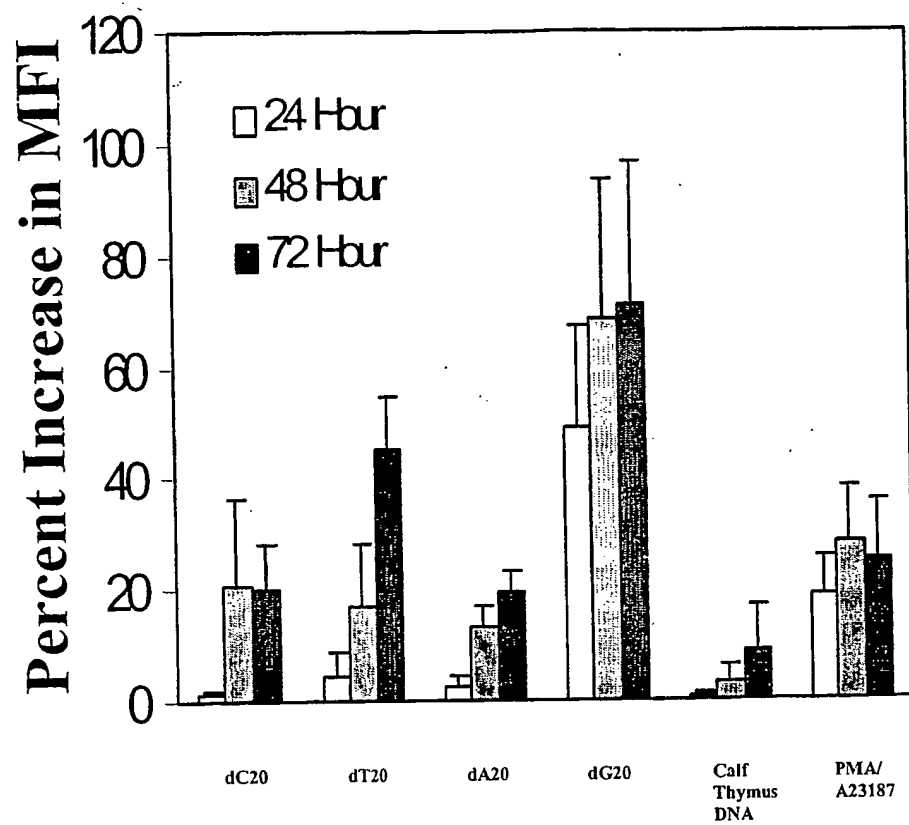
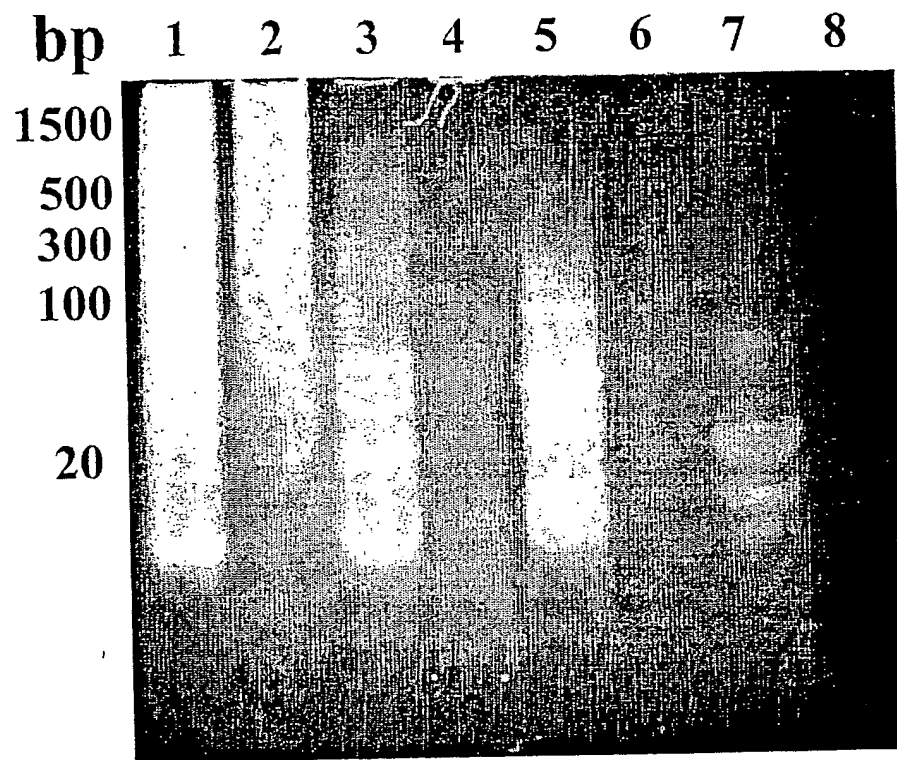
**Figure 7**

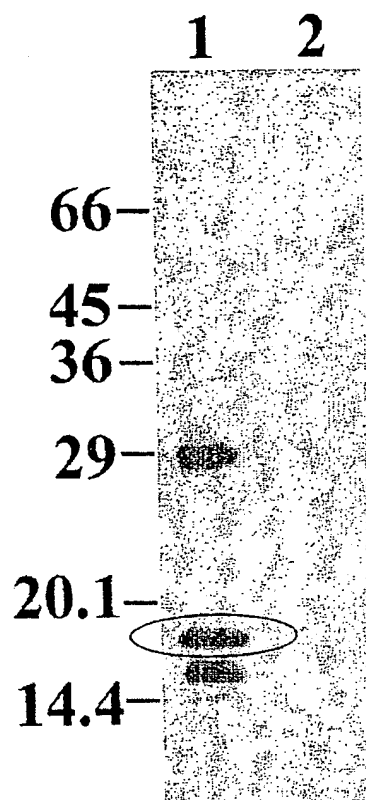
Figure 8



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Figure 9



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## Figure 10

```
1  CGGCACGAGGGTTCAATAGCATCTCAAGGCGCTTCAGAACTTAAAAGTTGA
    M S A Q A E E T A P E A A A P V 16
51  ACCATGTCGTCTCAGGCTGAGGAACTGCACCAGAAGCAGCAGCACCAGT
    Q P S Q P A A K K K G P A S K A 32
101 ACAACCATCACAACCAGCGGCCAAAAAGAAGGGACCCGCCAGTAAAGCAA
    K P A S A E K K N K K K G K G P 49
151 AGCCTGCCTCTGCAGAAAAAAGAACA AAAAGAAGAAAGGGAAAGGGCCC
    G K Y S Q L V I N A I Q T L G E R 66
201 GGAAAGTACAGCCAGCTGGTGATCAATGCTATCCAAACGCTGGGAGAGAG
    N G S S L F K I Y N E A K K V N 82
251 AAACGGCTCGTCTCTTTTAAAGATCTACAACGAGGCGAAGAAAGTGAAGT
    W F D Q Q H G R V Y L R Y S I R A 99
301 GGTTTGACCAGCAGCAGCGGGCGGTGTACCTCCGCTACTCCATCCGCGCG
    L L Q N D T L V Q V K G L G A N G 116
351 CTGCTGCAGAACGACACGCTCGTGCAGGTGAAGGGTCTGGGCGCCAACGG
    S F K L N K K K F I P R T K K S 132
401 CTCCTTCAAGCTCAACAAAAAGAAGTTCATCCCCAGAACCAAGAAGAGCT
    S V K P R K T A K P T K K P A K K 149
451 CTGTAAAGCCGAGAAAGACTGCGAAACCGACCAAAAAGCCAGCCAAAAAA
    A A K K K K R V S G V K K A T P P 166
501 GCAGCGAAGAAGAAGAAAAGGGTCAGCGGCGTGAAGAAGGCGACTCCCCC
    P E K T S K P K K A D K S P A V 182
551 CCCAGAGAAAACCTCCAACCCAAGAAAGCGGATAAAAGTCCAGCCGTCT
    S A K K A S K P K K A K Q T K K T 199
601 CTGCCAAGAAGGCGAGCAAGCCCAAGAAAGCTAAACAGACAAAAAAGACT
    A K K T * 203
651 GCTAAGAAGACTTAAAACGTTTATATTCTGCATGCTTTGTGCATTAAAGCA
701 TTGCACTGCGGGTAAACTGCACGCTTTCTGATCGCAGTTCATTAAGTAGG
751 ATATGCACAGTGTTTAAACCAAGTGTGCAAGTCACTCTGGTCTCAATGTTT
801 TACTGATGTAACCACATGTAAATAACTGTACAAAGAAGGAAACAATCACT
851 TTTGTAACGTCTGCTTTGTTATTATTTCTTTCTACTAGTTAGCTAAAAT
901 AACTGCTTATGGCTTCTTTTAAATATAATGATAAAAGAAAAAAAAAAAA
951 AAAAAA
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Figure 11

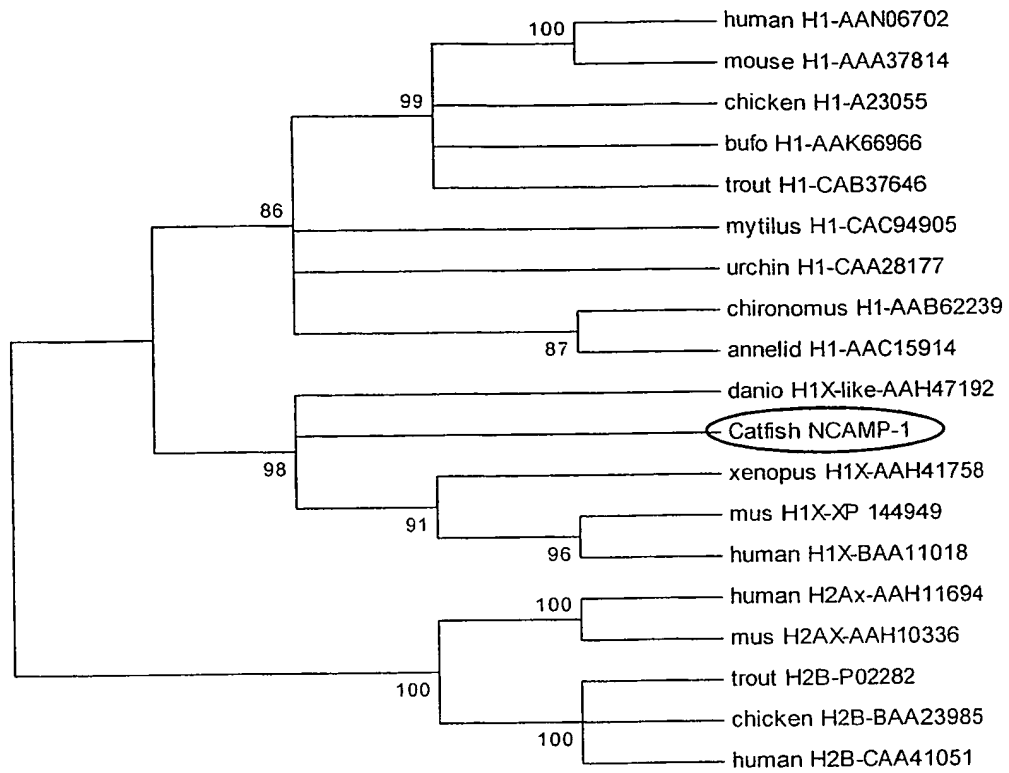
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Catfish NCAMP-1      : MSQAEEETAPRAAPVQPSQP-----AAKKGPASAKKPAEAKKNNKKKGKPGKYSQAVINATQTIGERNG : 68
Danio H1X-like-AAH47192 : -----MPAVVEESAPAPAP-----AEKAKAPAVAFSPAKK-----KKKSKGPGKYSKQVTDARTLGERNG : 59
Xenopus H1X-AAH41758 : -----MALELEENLHSTEEDEEEDEEDEMRSRSTRNKGGA-----SSSGNKKKKK--KKNQPGKYSQAVVDTRKLGERNG : 73
Mus H1X_XP_144949 : -----MSVELEEEALPPTSADG-----TAKTAKAGGSAAPTQPKRRKN--PKKNQPGKYSQAVVETIRKLGERRG : 63
human H1X-BAA11018 : -----MSVELEEEALPVTTAEG-----MAKKVTKAGGSAALSPSKKRNKSKKKNQPGKYSQAVVETIRRLGERNG : 64

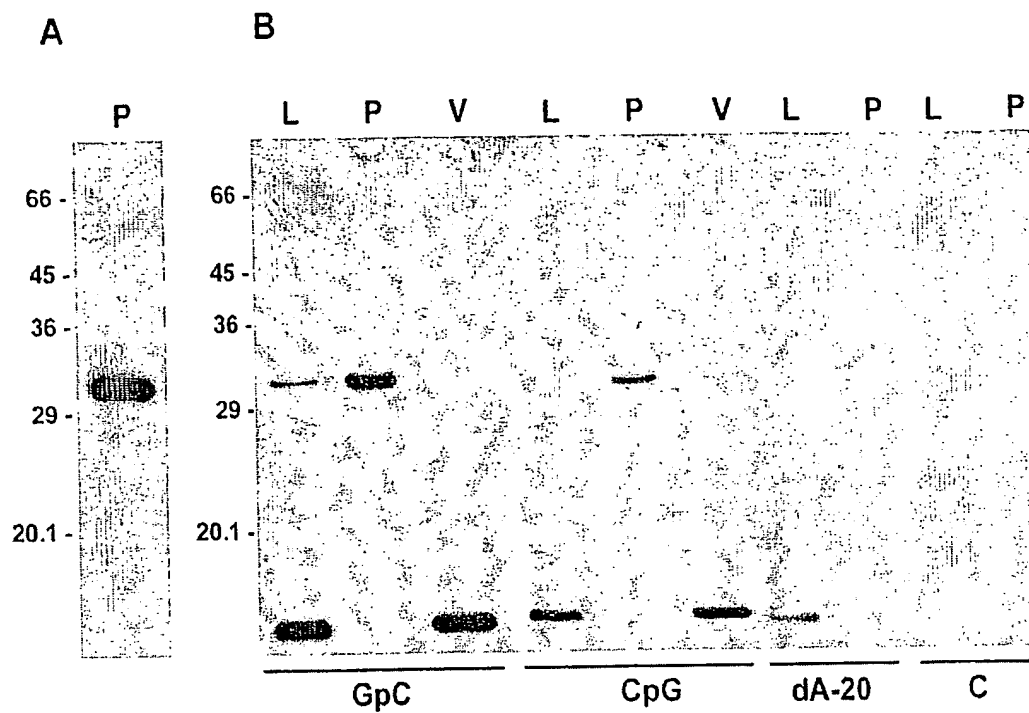
Catfish NCAMP-1      : SSLKIIYNEAKKVMFEDQOHGQGVYLRNSIRALVNDTLVQVKGCGANGSEKLNKKRFFIPRTKKSSVKPRKTAKTKKPAK : 148
Danio H1X-like-AAH47192 : SSLKIIYNEAKKVMFEDQKNGEMVILRSIRALVNDTLVQVKGCGANGSEKLNKKRLEKKPKK-AAKKATKKTETKPTSK : 138
Xenopus H1X-AAH41758 : SSLAKIYSEAKKVMFEDQONGRTYLYKSIKALVNDTLVQVKGCGANGSEFLNKKKLEGLPYDKPPAKPPSSSSSNKK : 153
Mus H1X_XP_144949 : SSLARIYAEARKVMFEDQONGRTYLYKSIKALVNDTLVQVKGCGANGSEFLNKKKLEGGAEERR-GASAASSPAPKAR-- : 140
human H1X-BAA11018 : SSLAKIYNEAKKVMFEDQONGRTYLYKSIKALVNDTLVQVKGCGANGSEFLNKKKLEGGERRGAPAAATAPAPTAKHA : 144

Catfish NCAMP-1      : KAAK-----KKRVSGVKNATPEPEKTSKPK-----KADKSFVSVSNKASPKPKAKQTKRTAKMT- : 203
Danio H1X-like-AAH47192 : KAVT-----KKVSAKKSARKSPVKKRTPKKT-----SVKKATAKPKTASAKPKPAAAKKTKSK-- : 192
Xenopus H1X-AAH41758 : QQQ-----GPFSSPSKSHKAKEPKAKAEKEKPKTSSAKAKSPKPKSAKKG-KRMKGAAPSVRPAKSKKA : 217
Mus H1X_XP_144949 : -----TAAADRTPAEPQ-PERRAHKS-----KKAAPASAKKVKVKAAPSPVPAKPKGK- : 188
human H1X-BAA11018 : KKAAPGAAGSRRADKKPAEGQKPEQSSHKKGAGAKKDKGKAKKTAAAGGKVKKPAAPSPVPAKPKGK- : 213

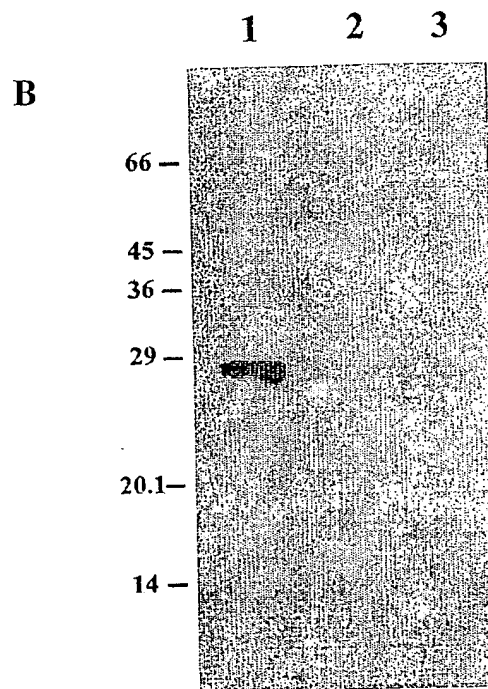
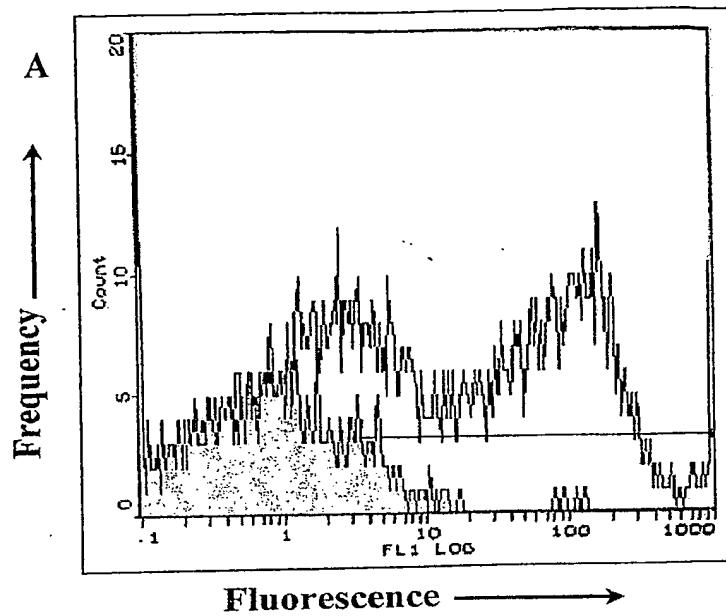
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**Figure 12**

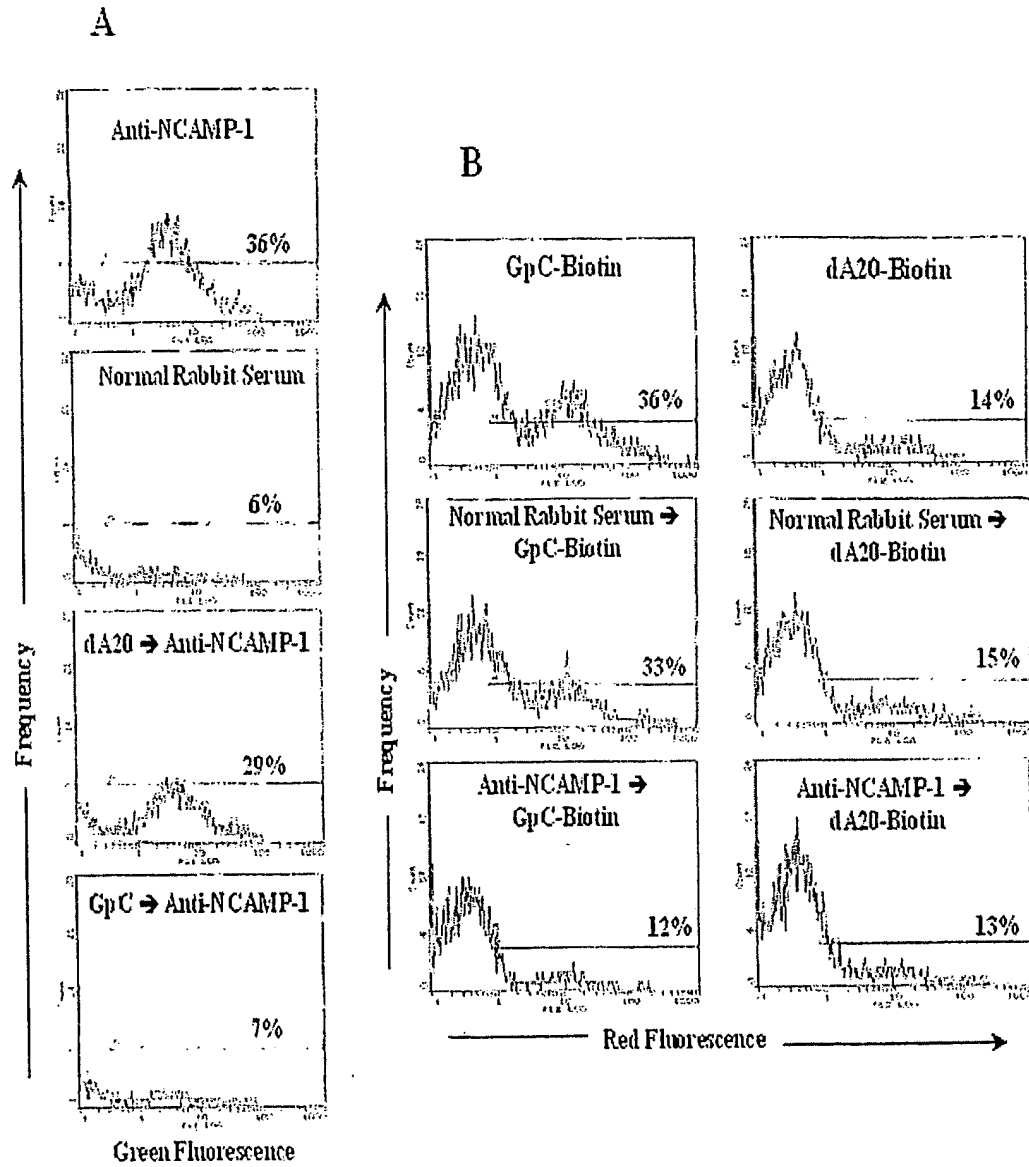
Figures 13A - 13B



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**Figures 14A - 14B****BEST AVAILABLE COPY**

## Figures 15A - 15B



## FIGURE 16

NCAMP-1      PEAAAPVQPSQPAAKKKGPASKAKPASAEKKNKKAAGKGPG  
 H1-Mus      -----SETAPAEKPPAKAE--  
 H1-Human      ---KLNKKAASGEAKPKAKAKSPKKAKA--  
 H1-Trout      -KAVAAKKSPPKAKKPAAT--

### C-Terminal residues :

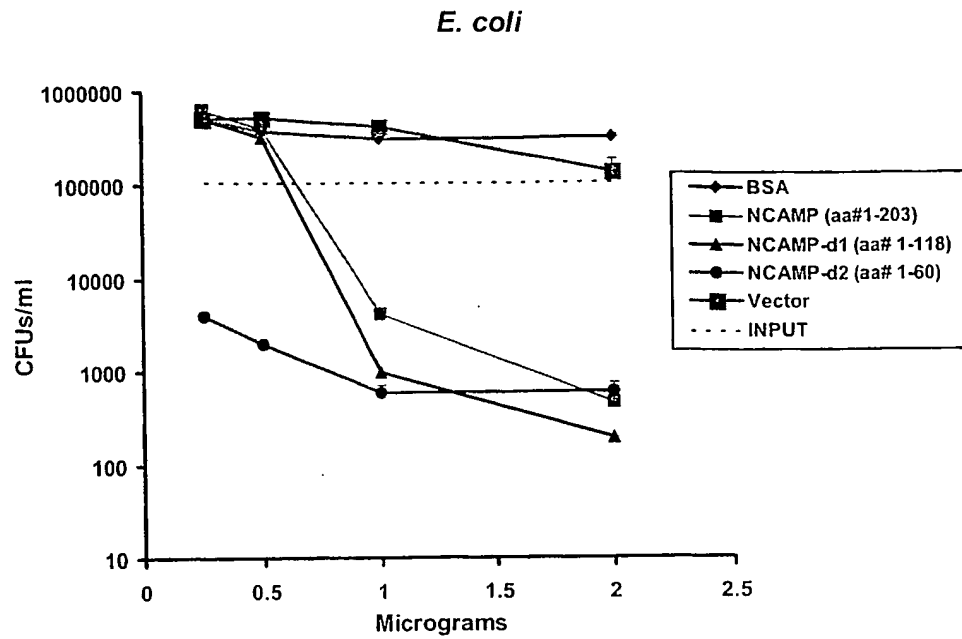
NCAMP-1  
 -TAKPTKKPAKKAACKKKRVSGVKKATPPPEKTSKPKKADKSPAVSAKKASKPKKAKQT  
 H2A CF --KGRGKQGKVRAKAKTRSS--  
 H2B Trout      -----EDEAKTAPKKGSKKAVTKXA--  
 H2B Bass1      -----PEPAKSAPKKGSKKAVT-  
 H2B Bass2      -----PDPAKPTAPKKGSKKAVTKTAG  
 H1-Trout      -----AEVAPAPAAAPAKAPKKKA  
 H1-Trout ---AEVAPAPAAAPAKAPKKKAAAKPKK-----

---



## Figures 17A - 17B

A



B

